

AMENDMENTS TO THE CLAIMS

This listing of claims will replace all prior versions and listings of all claims in the application.

Listing of Claims

Claims 1-59 (canceled)

60. (previously presented) A method of determining the identification of nucleotide(s) at a first detection position in a first domain of a target sequence, said target sequence comprising said first domain and a second domain, said method comprising:

- a. providing an electrode with a covalently attached capture probe, wherein said capture probe has a sequence substantially complementary to said second domain of said target sequence;
- b. contacting said electrode with:
 - (i) said target sequence;
 - (ii) a first label probe substantially complementary to said first domain, comprising a first nucleotide at an interrogation position and a first electron transfer moiety (ETM) with a first redox potential;
 - (iii) a second label probe substantially complementary to said first domain, comprising a second nucleotide at said interrogation position and a second ETM with a second redox potential;under conditions wherein if said nucleotide at said interrogation position is perfectly complementary to said detection position, hybridization of said probe(s) occurs; and
- c. detecting the presence of said first and/or second ETM to determine the nucleotide(s) at said detection position.

61. (previously presented) The method of Claim 60 wherein said method further comprises contacting said electrode with a third label probe substantially complementary to said first domain, comprising a third nucleotide at said interrogation position and a third ETM with a third redox potential.

62. (previously presented) The method of Claim 61 wherein said method further comprises contacting said electrode with a fourth label probe substantially complementary to said first domain, comprising a fourth nucleotide at said interrogation position and a fourth ETM with a fourth redox potential.
63. (previously presented) The method of Claim 60 wherein said electrode comprises an array of capture probes, each substantially complementary to a second domain of a different target sequence.
64. (previously presented) The method of Claim 60 wherein said first label probe contains a plurality of first ETMs.
65. (previously presented) The method of Claim 60 wherein said second label probe contains a plurality of second ETMs.
66. (previously presented) The method of Claim 60 wherein said electron transfer moieties comprise a transition metal complex.
67. (previously presented) The method of Claim 66 wherein said transition metal complex comprises a metallocene.
68. (previously presented) The method of Claim 67 wherein said metallocene is a ferrocene.
69. (previously presented) The method of Claim 67 wherein said metallocene is a ferrocene derivative.